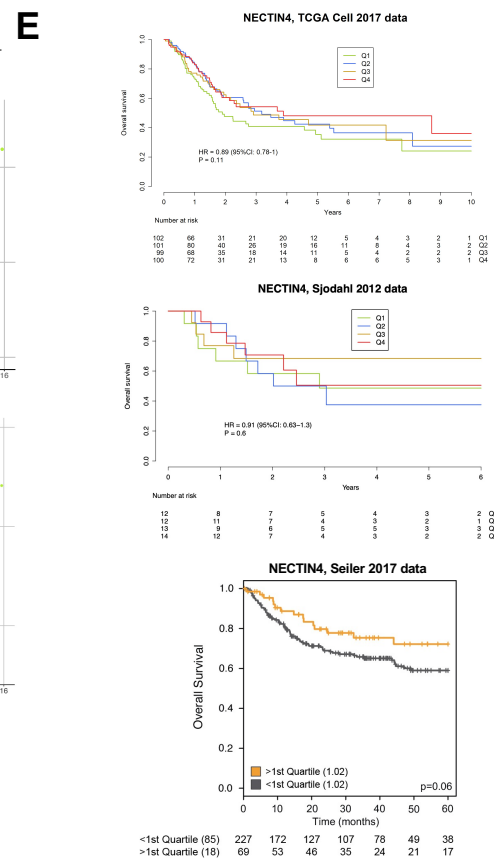


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Gene name Entrez ID	LumP	LumNS	LumU	Stroma- rich	Ba/Sq	NE-like
<i>NECTIN4</i> 81607	1.02	1.07	1.41	0.54	0.47	-0.44
<i>FOXA1</i> 3169	1.40	1.57	1.50	0.98	0.51	1.21
<i>GATA3</i> 2625	2.17	2.28	2.34	1.65	0.79	-0.35
<i>PPARG</i> 5468	1.16	1.05	1.32	0.48	0.34	-0.79



Supplementary Figure 2. Gene expression analysis of *NECTIN4* correlates. (A) Correlation between *NECTIN4* and *GATA3* mRNA expression across the Seiler 2017 (left), Sjødahl 2012 (middle), and TCGA (right) cohorts. (B) Correlation between *NECTIN4* and *FOXA1* mRNA expression across Seiler 2017 (left), Sjødahl 2012 (middle), and TCGA (right) cohorts. (C) Correlation between *NECTIN4* and *PPARG* mRNA expression across Seiler 2017 (left), Sjødahl 2012 (middle), and TCGA (right) cohorts. (D) Centroid values from the consensus molecular classifier by Kamoun et al. for *NECTIN4*, *FOXA1*, *GATA3*, and *PPARG*. (E) Kaplan-Meier curves of overall survival stratified by quartiles of *NECTIN4* mRNA expression in TCGA ($p=0.11$), Sjødahl 2012 ($p=0.6$), and Seiler 2017 ($p=0.06$) cohorts.